

PCT09

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/936,019

DATE: 10/04/2001
 TIME: 14:53:34

Input Set : A:\ES.txt
 Output Set: N:\CRF3\10042001\I936019.raw

3 <110> APPLICANT: Witcher, Derrick
 4 Rathnachalam, Radhakrishnan
 5 Micanovic, Radmila
 7 <120> TITLE OF INVENTION: Protease Resistant FLINT Analogs
 9 <130> FILE REFERENCE: X-13161
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/936,019
 C--> 12 <141> CURRENT FILING DATE: 2001-09-07
 14 <160> NUMBER OF SEQ ID NOS: 8
 16 <170> SOFTWARE: PatentIn Ver. 2.0
 18 <210> SEQ ID NO: 1
 19 <211> LENGTH: 271
 20 <212> TYPE: PRT
 21 <213> ORGANISM: Homo sapiens
 23 <400> SEQUENCE: 1
 25 Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu
 26 1 5 10 15
 28 Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro
 29 20 25 30
 31 Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His
 32 35 40 45
 34 Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val
 35 50 55 60
 37 Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His
 38 65 70 75 80
 40 Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe
 41 85 90 95
 43 Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro
 44 100 105 110
 46 Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr
 47 115 120 125
 49 Phe Ser Ala Ser Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn
 50 130 135 140
 52 Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His
 53 145 150 155 160
 55 Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val
 56 165 170 175
 58 Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe
 59 180 185 190
 61 Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu
 62 195 200 205
 64 Ala Pro Glu Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu
 65 210 215 220
 67 Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp
 68 225 230 235 240
 70 Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met
 71 245 250 255
 73 Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His

ENTERED

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74          260          265          270
77 <210> SEQ ID NO: 2
78 <211> LENGTH: 813
79 <212> TYPE: DNA
80 <213> ORGANISM: Homo sapiens
82 <400> SEQUENCE: 2
83 gtggcagaaa caccaccta cccctggcgg gacgcagaga caggggagcg gctggtgtgc 60
84 gcccagtgcc cccaggcac ctttgtgcag cgcccggtgcc gccgagacag cccacgacg 120
85 tgtggcccgt gtccaccgcg ccactacacg cagttctgga actacctgga gcgctgccgc 180
86 tactgcaacg tcctctgcgg ggagcgtgag gaggaggcac gggcttgcca cgccaccac 240
87 aaccgtgcct gccgctgccg caccggcttc ttgcgcacg ctggtttctg cttggagcac 300
88 gcatcgtgtc cacctggtgc cggcgtgatt gccccgggca cccccagcca gaacacgcag 360
89 tgccagccgt gccccccagg caccttctca gccagcagct ccagctcaga gcagtgccag 420
90 cccacccgca actgcacggc cctgggcctg gccctcaatg tgccaggctc ttctcccat 480
91 gacaccctgt gcaccagctg cactggcttc cccctcagca ccagggtacc aggagctgag 540
92 gagtgtgagc gtgccgtcat cgactttgtg gctttccagg acatctccat caagaggctg 600
93 cagcggctgc tgcaggccct cgaggccccg gagggtgagg gtccgacacc aaggcggggc 660
94 cgcgcggcct tgcagctgaa gctgcgtcgg cggtcacgg agctcctggg ggcgaggac 720
95 ggggcgctgc tggcgcggtc gctgcaggcg ctgcgcgtgg ccaggatgcc cgggctggag 780
96 cggagcgtcc gtgagcgctt cctccctgtg cac
97                                     813
98 <210> SEQ ID NO: 3
99 <211> LENGTH: 300
100 <212> TYPE: PRT
101 <213> ORGANISM: Homo sapiens
103 <400> SEQUENCE: 3
104 Met Arg Ala Leu Glu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val Leu
105   1          5          10          15
107 Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly Val Ala Glu
108   20          25          30
110 Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val
111   35          40          45
113 Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg
114   50          55          60
116 Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln
117   65          70          75          80
119 Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly
120   85          90          95
122 Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His Asn Arg Ala
123   100         105         110
125 Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu
126   115         120         125
128 His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro
129   130         135         140
131 Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala
132 145         150         155         160
134 Ser Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala
135   165         170         175
137 Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu
138   180         185         190

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140 Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala
141      195      200      205
143 Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile
144      210      215      220
146 Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu
147 225      230      235      240
149 Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys
150      245      250      255
152 Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu
153      260      265      270
155 Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu
156      275      280      285
158 Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His
159      290      295      300

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161 <210> SEQ ID NO: 4

162 <211> LENGTH: 29

163 <212> TYPE: PRT

164 <213> ORGANISM: Homo sapiens

166 <400> SEQUENCE: 4

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167 Met Arg Ala Leu Glu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val Leu
168 1      5      10      15

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170 Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly
171      20      25

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173 <210> SEQ ID NO: 5

174 <211> LENGTH: 39

175 <212> TYPE: DNA

176 <213> ORGANISM: Artificial Sequence

W--> 177 <220> FEATURE:

178 <223> OTHER INFORMATION: Description of Artificial Sequence: oligo primer

180 <400> SEQUENCE: 5

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181 gcaccagggt accaggagct gaggagtgtg agcgtgccg
182                                     39

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183 <210> SEQ ID NO: 6

184 <211> LENGTH: 44

185 <212> TYPE: DNA

186 <213> ORGANISM: Artificial Sequence

W--> 187 <220> FEATURE:

189 <223> OTHER INFORMATION: Description of Artificial Sequence: oligo primer

191 <400> SEQUENCE: 6

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192 tcagctgcaa ggcggcgcgccg cccgcttgtg gtgtcggacc ccag
193                                     44

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194 <210> SEQ ID NO: 7

195 <211> LENGTH: 44

196 <212> TYPE: DNA

197 <213> ORGANISM: Artificial Sequence

W--> 198 <220> FEATURE:

200 <223> OTHER INFORMATION: Description of Artificial Sequence: oligo primer

202 <400> SEQUENCE: 7

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203 ggggtccgac accacaagcg gggcgcgccg ccttgcagct gaag
204                                     44

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205 <210> SEQ ID NO: 8

206 <211> LENGTH: 43

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Input Set : A:\ES.txt
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207 <212> TYPE: DNA
208 <213> ORGANISM: Artificial Sequence
W--> 209 <220> FEATURE:
211 <223> OTHER INFORMATION: Description of Artificial Sequence: oligo primer
213 <400> SEQUENCE: 8
214 gcacagaatt catcagtgca cagggaggaa gcgctcacgg acg 43

VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:177 M:283 W: Missing Blank Line separator, <220> field identifier
L:187 M:283 W: Missing Blank Line separator, <220> field identifier
L:198 M:283 W: Missing Blank Line separator, <220> field identifier
L:209 M:283 W: Missing Blank Line separator, <220> field identifier